SEQUENCE LISTING

Warranger

Alland, David
Bloom, Barry R.
Jacobs Jr., William R.
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Val Val Pro Gly Leu Asn Leu Gly Ala Gly Asp Pro Met Ser Gly Leu 50 55 60

Arg Gln Ala Val Ala Ala Arg His Gly Phe Ala Gln Asp Val Ala Asn 65 70 75 . 80

Val Gly Phe Ala Gly Asp Ala Gly Ala Gly Val Ala Ser Val Ile Thr 85 90 95

Thr Asp Val Gly Ala Gly Leu Ala Ser Gly Leu Gly Ala Gly Phe Leu 100 105 110

Gly Gln Gry Gly Leu Ala Leu Ala Ala Ser Ser Gly Gly Phe Gly Gly Gln Val Gly Leu Ala Ala Gln Val Gly Leu Gly Phe Thr Ala Val Ile Glu Ala Glu Val Gly Ala Gln Val Gly Ala Gly Leu Gly Ile Gly Thr Gly Leu Gly Ala Gln Ala Gly Met Gly Phe Gly Gly Gly Val Gly Leu Gly Leu Gly Gly Gln Ala Gly Gly Val Ile Gly Gly Ser Ala Ala Gly Ala Ile Gly Ala Gly Val Gly Gly Arg Leu Gly Gly Asn Gly Gln Ile Gly Val Ala Gly Gln Gly Ala Val Gly Ala Gly Val Gly Ala Gly Val Gly Gly Gln Ala Gly Ile Ala Ser Gln Ile Gly Val Ser Ala Gly Gly Gly Leu Gly Gly Val Gly Asn Val Ser Gly Leu Thr Gly Val Ser Ser Asn Ala Val Leu Ala Ser Asn Ala Ser Gly Gln Ala Gly Leu Ile Ala Ser Glu Gly Ala Ala Leu Asn Gly Ala Ala Met Pro His Leu Ser Gly Pro Leu Ala Gly Val Gly Val Gly Gln Ala Gly Ala Ala Gly Gly Ala Gly Leu Gly Phe Gly Ala Val Gly His Pro Thr Pro Gln Pro Ala

13.5 ×

Val Val Gly Gly Val Gly Gly Ala Thr Ala Ala Gly Val Gly Gly Ala 340 345 350

Ala Leu Gly Ala Ala Gly Val Val Ala Lys Thr Glu Ala Ala Ala Gly

His Gly Asp Ile Leu Gly His Glu Gly Ala Ala Leu Gly Ser Val Asp 355 360 365

Thr Val Asn Ala Gly Val Thr Pro Val Glu His Gly Leu Val Leu Pro Ser Gly Pro Leu Ile His Gly Gly Thr Gly Gly Tyr Gly Gly Met Asn Pro Pro Val Thr Asp Ala Pro Ala Pro Gln Val Pro Ala Arg Ala Gln Pro Met Thr Thr Ala Ala Glu His Thr Pro Ala Val Thr Gln Pro Gln His Thr Pro Val Glu Pro Pro Val His Asp Lys Pro Pro Ser His Ser Val Phe Asp Val Gly His Glu Pro Pro Val Thr His Thr Pro Pro Ala Pro Ile Glu Leu Pro Ser Tyr Gly Leu Phe Gly Leu Pro Gly Phe <210> 4 <211> 640 <212> PRT <213> Mycobacterium tuberculosis Met Val Pro Ala Gly Leu Cys Ala Tyr Arg Asp Leu Arg Arg Lys Arg Ala Arg Lys Trp Gly Asp Thr Val Thr Gln Pro Asp Asp Pro Arg Arg Val Gly Val Ile Val Glu Leu Ile Asp His Thr Ile Ala Ile Ala Lys Leu Asn Glu Arg Gly Asp Leu Val Gln Arg Leu Thr Arg Ala Arg Gln Arg Ile Thr Asp Pro Gln Val Arg Val Val Ile Ala Gly Leu Leu Lys Gln Gly Lys Ser Gln Leu Leu Asn Ser Leu Leu Asn Leu Pro Ala Ala

Arg Val Gly Asp Asp Glu Ala Thr Val Val Ile Thr Val Val Ser Tyr \$100\$ \$100\$

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Gly	Ala	Pro	Ser	Pro 165	Leu	Leu	Arg	Gly	Gly 170	Leu	Ala	Phe	Ile	Asp 175	Thr
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Glu	Phe 210	Thr	Glu	Pro	Glu	Met 215	Trp	Phe	Val	Arg	Gln 220	Ala	His	Gln	Ile
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Trp	Arg	Glu	Ile	Val 245	Asn	Ala	Asn	Ala	Ala 250	His	Leu	Gln	Arg	Ala 255	Arg
Val	Pro	Met	Pro 260	Ile	Ile	Ala	Val	Ser 265	Ser	Leu	Leu	Arg	Ser 270	His	Ala
Val	Thr	Leu 275	Asn	Asp	Lys	Glu	Leu 280	Asn	Glu	Glu	Ser	Asn 285	Phe	Pro	Ala
Ile	Val 290	Lys	Phe	Leu	Ser	Glu 295	Gln	Val	Leu	Ser	Arg 300	Ala	Thr	Glu	Arg
Val 305	Arg	Ala	Gly	Val	Leu 310	Gly	Glu	Ile	Arg	Ser 315	Ala	Thr	Glu	Gln	Leu 320
Ala	Val	Ser	Leu	Gly 325	Ser	Glu	Leu	Ser	Val 330	Val	Asn	Asp	Pro	Asn 335	Leu
Arg	Asp ·	Arg	Leu 340	Ala	Ser	Asp	Leu	Glu 345	Arg	Arg	Lys	Arg	Glu 350	Ala	Gln
Gln	Ala	Val	Gln	Gln	Thr	Ala	Leu	Trp	Gln	Gln	Val	Leu	Gly	Asp	Gly

rne	370	ASP	Leu	Int	Ala	375	Val	Ash	urs	АЗР	380	Arg	Int	Arg	rne
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His	Lys	Met	Ile	Ile 485	Gly	Met	Arg	Gly	Ser 490	Tyr	Gly	Gly	Val	Val 495	Met
Ile	Gly	Met	Leu 500	Ser	Ser	Val	Val	Gly 505	Leu	Gly	Leu	Phe	Asn 510	Pro	Leu
Ser	Val	Gly 515	Ala	Gly	Leu	Ile	Leu 520	Gly	Arg	Met	Ala	Tyr 525	Lys	Glu	Asp
Lyš	Gln 530	Asn	Arg	Leu	Leu	Arg 535	Val	Arg	Ser	Glu	Ala 540	Lys	Ala	Asn	Val
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Asp	Arg	Leu	Lys	Met 565	Ile	Gln	Arg	Leu	Leu 570	Arg	Asp	His	Tyr	Arg 575	Glu
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Leu Ala Gly Thr Leu Lys Ala Gly Lys Ser Thr Leu Val Asn Ala Leu 50 55 60

Val Gly Asp Asp Ile Ala Pro Thr Asp Ala Thr Glu Ala Thr Arg Ile
65 70 75 80

Val Thr Trp Phe Arg His Gly Pro Thr Pro Arg Val Thr Ala Asn His 85 90 95

Arg Gly Gly Arg Arg Ala Asn Val Pro Ile Thr Arg Arg Gly Gly Leu 100 105 110

Ser Phe Asp Leu Arg Arg Ile Asn Pro Ala Glu Leu Ile Asp Leu Glu 115 120 125

Val Glu Trp Pro Ala Glu Glu Leu Ile Asp Ala Thr Ile Val Asp Thr 130 135 140

Pro Gly Thr Ser Ser Leu Ala Cys Asp Ala Ser Glu Arg Thr Leu Arg 145 150 155 160

Leu Leu Val Pro Ala Asp Gly Val Pro Arg Val Asp Ala Val Phe 165 170 175

Leu Leu Arg Thr Leu Asn Ala Ala Asp Val Ala Leu Leu Lys Gln Ile

Gly Gly Leu Val Gly Gly Ser Val Gly Ala Leu Gly Ile Ile Gly Val Ala Ser Arg Ala Asp Glu Ile Gly Ala Gly Arg Ile Asp Ala Met Leu Ser Ala Asn Asp Val Ala Lys Arg Phe Thr Arg Glu Leu Asn Gln Met Gly Ile Cys Gln Ala Val Val Pro Val Ser Gly Leu Leu Ala Leu Thr Ala Arg Thr Leu Arg Gln Thr Glu Phe Ile Ala Leu Arg Lys Leu Ala Gly Ala Glu Arg Thr Glu Leu Asn Arg Ala Leu Leu Ser Val Asp Arg Phe Val Arg Arg Asp Ser Pro Leu Pro Val Asp Ala Gly Ile Arg Ala . 300 Gln Leu Leu Glu Arg Phe Gly Met Phe Gly Ile Arg Met Ser Ile Ala Val Leu Ala Ala Gly Val Thr Asp Ser Thr Gly Leu Ala Ala Glu Leu Leu Glu Arg Ser Gly Leu Val Ala Leu Arg Asn Val Ile Asp Gln Gln Phe Ala Gln Arg Ser Asp Met Leu Lys Ala His Thr Ala Leu Val Ser Leu Arg Arg Phe Val Gln Thr His Pro Val Pro Ala Thr Pro Tyr Val Ile Ala Asp Ile Asp Pro Leu Leu Ala Asp Thr His Ala Phe Glu Glu Leu Arg Met Leu Ser Leu Leu Pro Ser Arg Ala Thr Thr Leu Asn Asp Asp Glu Ile Ala Ser Leu Arg Arg Ile Ile Gly Gly Ser Gly Thr Ser

...

Ala Ala Arg Leu Gly Leu Asp Pro Ala Asn Ser Arg Glu Ala Pro

435 440 445

Arg Ala Ala Leu Ala Ala Gln His Trp Arg Arg Ala Ala His 450 455 Pro Leu Asn Asp Pro Phe Thr Thr Arg Ala Cys Arg Ala Ala Val Arg 465 470 475 Ser Ala Glu Ala Met Val Ala Glu Phe Ser Ala Arg Arg 485 490 <210> 6 <211> 25 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Uniamp primer sequence <400> 6 cctctgaagg ttccagaatc gatag 25 <210> 7 <211> 32 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Uniamp XhoI adapter sequence top strand <400> 7 cctctgaagg ttccagaatc gatagctcga gt 32 <210> 8 <211> 35 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Uniamp XhoI adapter sequence bottom strand

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